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Nucleotide Sequence									
1	10	20	30	40	50	60	70	80	
1	GGACGTCGAG	GCATTACANT	CGCGAATCCA	AGCCNTAGCA	TGAAACAGCG	AGCITGACGC	CTCACCGACG	AGTCTCAACT	80
81	AAAGGGACT	CCCGGAGCTA	GGGTGGGA	CTCGGCTCA	CACAGTGAGT	GCCGGCTATT	GGACTTTTGT	CCAGTGACAG	160
161	CTGAGACAC	AAGGACACG	GGAGGAGTG	TAGGAGAGAA	CGCCCGGNA	CAGCGATCGC	CCAGCACCAA	GTCCGCTTCC	240
241	AGGCTTTCG	TTTCTTTGCC	TCCATCTTGG	GTCCGCCCTTC	CCGGCGTCTA	GGGAGCGNA	GGCTGAGGTG	GCAGCGGCAG	320
321	GAGAGTCCG	CCCGACAGG	ACGACTCCC	CCACTGGAAA	GGATTCTGNA	AGAAATGAAG	TCAGCCCTCA	GAATGNAAT	400
401	TGACTGCCTG	CTGCTTTCC	TGTTGACTGG	CCCGGAGCTG	TACTGCAAGA	CCCTTGTGAG	CTTCCCTTAGT	CTAAGAGTAG	480
481	GATGCTGCT	GAACTCATCC	NTCAGGTTGA	AGNAGCACTT	GATACAGATG	AGAGGAGAT	GCTGCTCTTT	TTGTGCCGGG	560
561	ATGTTGCTAT	AGATGTGGTT	CCACCTAATG	TCAGGGACCT	TCTGGTATTT	TTACGGGAAA	GAGGTAACT	GTCTGTCGGG	640
641	GACTTGGCTG	AATGCTCTA	CAGAGTGAGG	CGATTTGACC	TGCTCNAACG	TATCTTGAG	ATGGACAGAA	AAGCTGTGGA	720
721	GACCCACCTG	CTCAGGAAAC	CTCACCITGT	TTCGGACTAT	AGAGTGCTGA	TGGCAGAGAT	TGGTGAGGNT	TTGGATTAAT	800
801	CTGATGTGTC	CTCATTAAT	TTCCTCANTG	AGGATTAACN	GGCCCGAGGC	AAGNTAAGCA	AGGAGAAGNG	TTTCTTGGAC	880
881	CTGTGGTTG	AGTTGGAGAA	ACTAATTTG	GTTGCCCCAG	ATCMACTGGA	TTTATTAAGAA	AATGCCCTAA	AGAACNTCCA	960
961	CAGAAATAG	CTGAAGACNA	AAATCCAGAA	GTACAAAGCAG	TCTGTTCAG	GAGCAGGGAC	AAGTTACAGG	AATGTTCTCC	1040
1041	AAGCAGCAAT	CCAAAGAGT	CTCAGGATC	CTTCANATTA	CTTCAGGATG	ATAACACCCCT	ATGCCCATTG	TCCTGATCTG	1120
1121	AAATTCCTG	GAATTTGTT	CATGAGNTTA	ACNTGGNACT	GCCCTACTACT	AATCAATCTG	AATGATTAAA	TCGTTTTCAT	1200
1201	TTCTAAATGT	GTATATATGT	GTTAGCCCT	TTCTTGTGTC	TGTATGTTTA	GATGCTTTCC	AATCTTTTGT	TACTACTAAT	1280
1281	AATGCTATAA	AATAANTATC	CTTGTACTTC	TTAANAANA	AAAAAAAAAA	AAAAAAAAAA	AAAAAAAAAA	AAAAAAAAAA	1360
1361	AAAAAAAAAA	AAA							1373

Deduced Amino Acid Sequence

MSAEVINQVEEALDTDEKEMLLFLCRDVAIDVPPNVRLLDILRERKLSVGDLAELLYRVRRFDLLKRILKMDRKAVE
 THLLRNPHLVSDYRVLMAEI GEDLDKSDVSSLIFLMKDYMGRIKISKEKSFDLVVELEKLNLVAPDQLDLEKCLKNIIH
 RIDLKTAKIQKYKQSVQAGTSYRNVLAQAIQSLKDPNNFRMITPYAHCPDLKILGNCSEWZ

FIG. 2

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Deduced Amino Acid Sequence

MSAEVTHQVEEAEEDTDEKEMLLFLCRDVAIDVPPMVRDLEDEPERERGKLSVGDIAEELLYRARRFELKKKTEKMDKAVE
 TMLRNPHLVSDYRVLMAEIGEDLDKSDVSSLIIFLMKDYMGKGKISKESFLDLVVELEKLNLVAPDQDLLEKCLKNIH
 RIDLTKIQKYKQSVQAGTSYRNVLQAAIQKSLKDPSSNFRLNHNGRSKEQRLKEQLGAQQEPVKKSIQSEAFLPQSIP
 EERYKMKSKPLGICLIIDCIGNETELLRTFTSLGYEVQKFLHLSMHGISQILGQFACMPEHRDYDSFVCVLVSRGGSQS
 VYGVDDQTHSGLPLHHIRRMFGDSCPYLAGPKMFFIQNYVVSEGQLENSLLLEVDPAMKNVEFKAQKRGCLCTVHREAD
 FFWSLCTADMSLLEQSHSSPSLYLQCLSQKLRQERKRPLLDLHIELNGYMYDWNRSRVSAKEYVYVWLQHTLRKKLILSYT

FIG. 1B

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mCASHa	DITQIVARYASMAQ	HQDYDSFA	CVLVSL	CGSQSMGRDQVMSGFS	LDHVKNMFT	GDTCPSLR	GKPKL	FFIQNYESLGS	370
NCASHa	GISQILGQFACMPE	HROYDSFV	CVLVSL	SRGGSQSVYGVQTHSGLP	LHHIRRMFM	GDSCPV	LAGKPKM	FFIQNYVYVSEG	385
CASP-8	EQIYEIKIYQLMD	HSNMDCFC	ICILSHGDKG	IYGTIDGQEG	PIYELT	SOFTGL	KGPSLAGKPKV	FFIQACGGDN	384
CASP-10	EMEMVLQKQKCNPA	HADGDC	FVFCILTHGRFGAVYS	DEALI	PIREIMSH	FITALQCP	RLAEKPKL	FFIQACGGGEE	362
CASP-3	EELVELMRDVSKE	HSKRSS	FV	CVLLSHGEEQ	IIFGTNG	PV	DLKKITN	FERGDRCS	166
CASP-1	SDMTTELEAF	AHRPE	HKTS	DSSTFLVFM	SHGIREGIC	CGKKHSEQ	VPD	ILQLNAL	IFNMLNTKNCPS	288
mCASHa	QLEDSS	LEVDGPS	IKNVDSKPLQPR	HCITTH	FEADL	FWSLCT	ADVSHLEK	PSSSSSV	442
NCASHa	QLENSLL	LEVDGPAMK	NVEFKAQKRG	CTVHRE	ADFFWSL	CTADMSL	LEQSHSS	SPSLV	438
CASP-8	YQKGIP	VEIDSEEQ	PYLEMDLSSP	QTRYIP	DEADFL	LLGMAT	IVNNCV	SYRNP	435
CASP-10	IQPSVS	IEADALN	PEQAPTS	LQDSIPA	EADFL	LLGLAT	IVPGYVS	429
CASP-3	LOC	GIETDSGVDD	MACHKIPV	EADFL	LYAYS	IAPGYYS	427
CASP-1	PGVVWF	KD	SVGVSGNLS	LPTEEFED	AIKKAH	IERKDF	IAFCSS	381
mCASHa	LVDLHVELMOK	VYAWN	SGVSSKEKYS	LSLQ	HTLR	KKLL	ILAPT	486
NCASHa	LVDLHIEL	NGY	MYDWN	SRVSAKEKY	YVWLQ	HTLR	KKLL	LSYT	481
CASP-8	GDILTL	LTETV	NYEV	SNKDDKKNMG	KMPQPT	FTLR	KKLV	FFPSD	479
CASP-10	HEILSL	ILTAV	NDV	SRVVDKQG	TKKMPQPA	FTLR	KKLV	FFVPLDALS	480
CASP-3	LEFMH	ILTRV	NRKV	ATEFES	FSFDT	FAHKK	QIIP	CIV	SM	277
CASP-1	COVEE	IFAKV	RF	SFEQ	PDGRA	QMD	TT	TERV	404

FIG. 3C